CELL. cann: JENSEN, M. i: April 30, 2001 Examiner: Unassigned Our Docket No.: 24751-2502

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Serial No.: 09/846,637 Art Unit: 1645

+3 MAD CACCGGCGAA GGAGGATCGA ATTCCTGCAG CCCGCTATCT GCAGGCCGCC ACCATGGCCG 1 GTGGCCGCTT CCTCCTAGCT TAAGGACGTC GGGCGATAGA CGTCCGGCGG TGGTACCGGC +3 DYLI G G T S Y V P D D S G L T ACTACCTGAT TAGTGGGGGC ACGTCCTACG TGCCAGACGA CGGACTCACA GCACAGCAGC TGATGGACTA ATCACCCCCG TGCAGGATGC ACGGTCTGCT GCCTGAGTGT CGTGTCGTCG +3 L F N C G D G L T Y N D F L I L PGYID TCTTCAACTG CGGAGACGGC CTCACCTACA ATGACTTTCT CATTCTCCCT GGGTACATCG 121 AGAAGTTGAC GCCTCTGCCG GAGTGGATGT TACTGAAAGA GTAAGAGGGA CCCATGTAGC D Q V +3 D L T SALT KKI TLKT ACTTCACTGC AGACCAGGTG GACCTGACTT CTGCTCTGAC CAAGAAAATC ACTCTTAAGA 181 TGAAGTGACG TCTGGTCCAC CTGGACTGAA GACGAGACTG GTTCTTTTAG TGAGAATTCT T-P-L-T-S-S-T-M-B-T-V-T-S-A-G-M-A CCCCACTGGT TTCCTCTCC ATGGACACAG TCACAGAGGC TGGGATGGCC ATAGCAATGG GGGGTGACCA AAGGAGAGGG TACCTGTGTC AGTGTCTCCG ACCCTACCGG TATCGTTACC +3 ALTG G I G F I H H N C T Р E F CGCTTACAGG CGGTATTGGC TTCATCCACC ACAACTGTAC ACCTGAATTC CAGGCCAATG 301 GCGAATGTCC GCCATAACCG AAGTAGGTGG TGTTGACATG TGGACTTAAG GTCCGGTTAC Y E Q +3 V R K V K K G F I T D P V 361 AAGTTCGGAA AGTGAAGAAA TATGAACAGG GATTCATCAC AGACCCTGTG GTCCTCAGCC TTCAAGCCTT TCACTTCTTT ATACTTGTCC CTAAGTAGTG TCTGGGACAC CAGGAGTCGG +3 K D R V R D V F E Α KAR H G F CGIP CCAAGGATCG CGTGCGGGAT GTTTTTGAGG CCAAGGCCCG GCATGGTTTC TGCGGTATCC 421 GGTTCCTAGC GCACGCCCTA CAAAAACTCC GGTTCCGGGC CGTACCAAAG ACGCCATAGG +3 T G R M G S R L V G I I S SRDI CAATCACAGA CACAGGCCGG ATGGGGAGCC GCTTGGTGGG CATCATCTCC TCCAGGGACA 481 GTTAGTGTCT GTGTCCGGCC TACCCCTCGG CGAACCACCC GTAGTAGAGG AGGTCCCTGT IDFLKEE E H D C F L E E I M +3 TKRE TTGATTTTCT CAAAGAGGAG GAACATGACT GTTTCTTGGA AGAGATAATG ACAAAGAGGG 541 AACTAAAAGA GTTTCTCCTC CTTGTACTGA CAAAGAACCT TCTCTATTAC TGTTTCTCCC +3 EDLVVAP A G I T L K E A N E AAGACTTGGT GGTAGCCCCT GCAGGCATCA CACTGAAGGA GGCAAATGAA ATTCTGCAGC 60i TTCTGAACCA CCATCGGGGA CGTCCGTAGT GTGACTTCCT CCGTTTACTT TAAGACGTCG +3 SKKGKL P I V NEDD ELV 661 GCAGCAAGAA GGGAAAGTTG CCCATTGTAA ATGAAGATGA TGAGCTTGTG GCCATCATTG CGTCGTTCTT CCCTTTCAAC GGGTAACATT TACTTCTACT ACTCGAACAC CGGTAGTAAC T D +3 L K K N R D Y P L A S K D 721 CCCGGACAGA CCTGAACAAC AATCCGGACT ACCCACTAGC CTCCAAAGAT GCCAAGAAAC GGGCCTGTCT GGACTTCTTC TTAGCCCTGA TGGGTGATCG GAGGTTTCTA CGGTTCTTTG +3G Α Α I G T H E D D K Y R AGCTGCTGTG TGGGGCAGCC ATTGGCACTC ATGAGGATGA CAAGTATAGG CTGGACTTGC 781 TCGACGACAC ACCCCGTCGG TAACCGTGAG TACTCCTACT GTTCATATCC GACCTGAACG +3 LAQA G V D V V V L D S S QGN S I F Q TCGCCCAGGC TGGTGTGGAT GTAGTGGTTT TGGACTCTTC CCAGGGAAAT TCCATCTTCC 841 AGCGGGTCCG ACCACACCTA CATCACCAAA ACCTGAGAAG GGTCCCTTTA AGGTAGAAGG I K Y I K D K Y P N L Q V I G G N +3 I N M AGATCAATAT GATCAAGTAC ATCAAAGACA AATACCCTAA TCTCCAAGTC ATTGGAGGCA 901 TCTAGTTATA CTAGTTCATG TAGTTTCTGT TTATGGGATT AGAGGTTCAG TAACCTCCGT

FIG. 1A

Title: SELECTION SYSTEM
CELL
Applicant: JENSEN, M.
Filed: April 30, 2001
Examiner: Unassigned
Our Docket No.: 24751-2502

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Serial No.: 09/846,637



+3 A A Q AKN LIDA GVD ALRV 961 ATGTGGTCAC TGCTGCCCAG GCCAAGAACC TCATTGATGC AGGTGTGGAT GCCCTGCGGG TACACCAGTG ACGACGGGTC CGGTTCTTGG AGTAACTACG TCCACACCTA CGGGACGCCC ICIIQEV VGMGSGS L A C +3 G R P Q 1021 TGGGCATGGG AAGTGGCTCC ATCTGCATTA TCCAGGAAGT GCTGGCCTGT GGGCGGCCCC ACCCGTACCC TTCACCGAGG TAGACGTAAT AGGTCCTTCA CGACCGGACA CCCGCCGGGG V Y K V Y E Y A R R F G V 1081 AAGCAACAGC AGTGTACAAG GTGTATGAGT ATGCACGGCG CTTTGGTGTT CCGGTCATTG TTCGTTGTCG TCACATGTTC CACATACTCA TACGTGCCGC GAAACCACAA GGCCAGTAAC VGHIAKA D G G I Q N L A L +3 CTGATGGAGG AATCCAAAAT GTGGGTCATA TTGCGAAAGC CTTGGCCCTT GGGGCCTCCA 1141 GACTACCTCC TTAGGTTTTA CACCCAGTAT AACGCTTTCG GAACCGGGAA CCCCGGAGGT T М Ε P М G S L L A A T G Ε Α 1201 CAGTCATGAT GGGCTCTCTC CTGGCTGCCA CCACTGAGGC CCCTGGTGAA TACTTCTTTT GTCAGTACTA CCCGAGAGAG GACCGACGGT GGTGACTCCG GGGACCACTT ATGAAGAAAA +3 R L K K Y R G M G S L D A CCGATGGGAT CCGCCTAAAG AAATATCGCG GTATGGGTTC TCTCGATGCC ATGGACAAGC 1261 GGCTACCCTA GGCCGATTTC TTTATAGCGC CATACCCAAG AGAGCTACGG TACCTGTTCG H L S S Q N R Y F S E A D K I K V A Q G V +3 ACCTCAGCAG CCAGAACAGA TATTTCAGTG AAGCTGACAA AATCAAAGTG GCCCAGGGAG 1321 TGGAGTCGTC GGTCTTGTCT ATAAAGTCAC TTCGACTGTT TTAGTTTCAC CGGGTCCCTC +3 V S G A V Q D K G S I H K F V P Y 1381 TGTCTGGTGC TGTGCAGGAC AAAGGGTCAA TCCACAAATT TGTCCCTTAC CTGATTGCTG ACAGACCACG ACACGTCCTG TTTCCCAGTT AGGTGTTTAA ACAGGGAATG GACTAACGAC GIQH SCQ DIG AKSL TQV 1441 GCATCCAACA CTCATGCCAG GACATTGGTG CCAAGAGCTT GACCCAAGTC CGAGCCATGA CGTAGGTTGT GAGTACGGTC CTGTAACCAC GGTTCTCGAA CTGGGTTCAG GCTCGGTACT Y S G E L K F E K R T S S A Q V +3 EGGV TGTACTCTGG GGAGCTTAAG TTTGAGAAGA GAACGTCCTC AGCCCAGGTG GAAGGTGGCG 1501 ACATGAGACC CCTCGAATTC AAACTCTTCT CTTGCAGGAG TCGGGTCCAC CTTCCACCGC +3 V H S L H S Y E K R L F 1561 TCCATAGCCT CCATTCGTAT GAGAAGCGGC TTTTCTGATC TAGCTCGACA TGATAAGATA AGGTATCGGA GGTAAGCATA CTCTTCGCCG AAAAGACTAG ATCGAGCTGT ACTATTCTAT CATTGATGAG TTTGGACAAA CCACAACTAG AATGCAGTGA AAAAAATGCT TTATTTGTGA 1621 GTAACTACTC AAACCTGTTT GGTGTTGATC TTACGTCACT TTTTTTACGA AATAAACACT AATTTGTGAT GCTATTGCTT TATTTGTGAA ATTTGTGATG CTATTGCTTT ATTTGTAACC 1681 TTAAACACTA CGATAACGAA ATAAACACTT TAAACACTAC GATAACGAAA TAAACATTGG ATTATAAGCT GCAATAAACA AGTTAACAAC AACAATTGCA TTCATTTTAT GTTTCAGGTT 1741 TAATATTCGA CGTTATTTGT TCAATTGTTG TTGTTAACGT AAGTAAAATA CAAAGTCCAA CAGGGGGAGG TGTGGGAGGT TTTTTAAAGC AAGTAAAACC TCTACAAATG TGGTAGATCA 1801 GTCCCCTCC ACACCCTCCA AAAAATTTCG TTCATTTTGG AGATGTTTAC ACCATCTAGT 1861 TTTAAATGTT AGCGAAGAAC ATGTGAGCAA AAGGCCAGCA AAAGGCCAGG AACCGTAAAA AAATTTACAA TCGCTTCTTG TACACTCGTT TTCCGGTCGT TTTCCGGTCC TTGGCATTTT AGGCCGCGTT GCTGGCGTTT TTCCATAGGC TCCGCCCCCC TGACGAGCAT CACAAAAATC TCCGGCGCAA CGACCGCAAA AAGGTATCCG AGGCGGGGG ACTGCTCGTA GTGTTTTTAG GACGCTCAAG TCAGAGGTGG CGAAACCCGA CAGGACTATA AAGATACCAG GCGTTTCCCC 1981 CTGCGAGTTC AGTCTCCACC GCTTTGGGCT GTCCTGATAT TTCTATGGTC CGCAAAGGGG

FIG. 1B

Sheet 3 of 7

Title: SELECTION SYSTEMS FOR GENETICALLY MODIFIED
CELL
ant: JENSEN, M.
April 30, 2001

Examiner: Unassigned
Our Docket No.: 24751-2502

1984657 USBAIR



2041	CTGGAAGCTC	CCTCGTGCGC	TCTCCTGTTC	CGACCCTGCC	GCTTACCGGA	TACCTGTCCG
	GACCTTCGAG	GGAGCACGCG	AGAGGACAAG	GCTGGGACGG	CGAATGGCCT	ATGGACAGGC
2101	CCTTTCTCCC	TTCGGGAAGC	GTGGCGCTTT	CTCAATGCTC	ACGCTGTAGG	TATCTCAGTT
	GGAAAGAGGG	AAGCCCTTCG	CACCGCGAAA	GAGTTACGAG	TGCGACATCC	ATAGAGTCAA
2161	CGGTGTAGGT	CGTTCGCTCC	AAGCTGGGCT	GTGTGCACGA	ACCCCCGTT	CAGCCCGACC
	GCCACATCCA	GCAAGCGAGG	TTCGACCCGA	CACACGTGCT	TGGGGGGCAA	GTCGGGCTGG
2221	GCTGCGCCTT	ATCCGGTAAC	TATCGTCTTG	AGTCCAACCC	GGTAAGACAC	GACTTATCGC
	CGACGCGGAA	TAGGCCATTG	ATAGCAGAAC	TCAGGTTGGG	CCATTCTGTG	CTGAATAGCG
2281	CACTGGCAGC	AGCCACTGGT	AACAGGATTA	GCAGAGCGAG	GTATGTAGGC	GGTGCTACAG
	GTGACCGTCG	TCGGTGACCA	TTGTCCTAAT	CGTCTCGCTC	CATACATCCG	CCACGATGTC
2341	AGTTCTTGAA	GTGGTGGCCT	AACTACGGCT	ACACTAGAAG	AACAGTATTT	GGTATCTGCG
		CACCACCGGA				
2401		GCCAGTTACC				
		CGGTCAATGG				
2461		TAGCGGTGGT				
		ATCGCCACCA				
2521		AGATCCTTTG				
		TCTAGGAAAC				
2581		GATTTTGGTC				
2001		CTAAAACCAG				
2641		GTGTGTTGGT				
2011		CACACAACCA				
2701		CAGGAAGGCA				
2/01		GTCCTTCCGT				
2761		ACAATCAACA				
2/61		TGTTAGTTGT				
2821		GGCCCGCCTG				
2021						
2001		CCGGGCGGAC				
2881		CCCATAGTAA				
0041		GGGTATCATT				
2941		ACTGCCCACT				
		TGACGGGTGA				
3001		AATGACGGTA				
		TTACTGCCAT				
3061		ACTTGGCAGT				
		TGAACCGTCA				
3121		TACATCAATG				
		ATGTAGTTAC				
3181		GACGTCAATG				
		CTGCAGTTAC				
3241		AACTCCGCCC				
		TTGAGGCGGG				
3301		AGAGCTCGTT				
		TCTCGAGCAA				
3361		CATAGAAGAC				
	AAAACTGGAG	GTATCTTCTG	TGGCCCTGGC	TAGGTCGGAG	GCGCCGGCCC	TTGCCACGTA

FIG. 1C

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3421					GCCTATAGAG	
	ACCTTGCGCC	TAAGGGGCAC	GGTTCTCACT	GCATTCATGG	CGGATATCTC	
3481		GCTTCTTATG		TGTTTTTGGC	TTGGGGTCTA	
					AACCCCAGAT	
3541		TTATAGGTGA			GTGTGGGTTA	TTGACCATTA
	GAAGGAGTAC	AATATCCACT			CACACCCAAT	
3601	TTGACCACTC	CCCTATTGGT	GACGATACTT	TCCATTACTA	ATCCATAACA	TGGCTCTTTG
	AACTGGTGAG	GGGATAACCA	CTGCTATGAA	AGGTAATGAT	TAGGTATTGT	ACCGAGAAAC
3661	CCACAACTCT	CTTTATTGGC	TATATGCCAA	TACACTGTCC	TTCAGAGACT	GACACGGACT
					AAGTCTCTGA	
3721	CTGTATTTTT	ACAGGATGGG	GTCTCATTTA	TTATTTACAA	ATTCACATAT	ACAACACCAC
		TGTCCTACCC				
 3781					ATCTCCACCC	
	GCAGGGGTCA	CGGGCGTCAA			TAGAGGTGCG	
3841		GGACATGGGC			GCTTCTACAT	
					CGAAGATGTA	
3901					TTGCTCCTAA	
	GAGGGTACGG	AGGTCGCTGA	GTACCAGCGA	GCCGTCGAGG	AACGAGGATT	GTCACCTCCG
3961					CCGCACAAGG	
					GGCGTGTTCC	
4021	AGGGTATGTG	TCTGAAAATG			ACCGCTGACG	
		AGACTTTTAC			TGGCGACTGC	GTAAACCTTC
4081	ACTTAAGGCA	GCGGCAGAAG			GTTGTGTTCT	
					CAACACAAGA	
4141	AGAGGTAACT	CCCGTTGCGG	TGCTGTTAAC	GGTGGAGGGC	AGTGTAGTCT	GAGCAGTACT
					TCACATCAGA	
4201					ACTAACAGAC	
	GCAACGACGG	CGCGCGCGGT	GGTCTGTATT	ATCGACTGTC	TGATTGTCTG	ACAAGGAAAG
				MCS		
					*****	
4261					GTAGCTCTAG	
					CATCGAGATC	
4321					TTTGTCTATA	
					AAACAGATAT	ACAATAAAAG
4381		CCGTCTTTTG				TCTTCTTGAC
					GGACCGGGAC	
4441		AGGGGTCTTT				TGAATGTCGT
					GTTCCAGACA	
4501					ACGTCTGTAG	
					TGCAGACATC	
4561					GGCCAAAAGC	
					CCGGTTTTCG	
4621					GTGAGTTGGA	
					CACTCAACCT	
4681					CTGAAGGATG	
					GACTTCCTAC	
4741					TGCTTTACAT	
	TGGGGTAACA	TACCCTAGAC	TAGACCCCGG	AGCCACGTGT	ACGAAATGTA	CACAAATCAG

FIG. 1D

4801	GAGGTTAAAA	AAACGTCTAG	GCCCCCGAA	CCACGGGGAC	GTGGTTTTCC	TTTGAAAAAC
	CTCCAATTTT	TTTGCAGATC	CGGGGGGCTT	GGTGCCCCTG	CACCAAAAGG	AAACTTTTTG
4861	ACGATAATAC	CATGGGTAAG	TGATATCTAC	TAGTTGTGAC	CGGCGCCTAG	TGTTGACAAT
	TGCTATTATG	GTACCCATTC	ATCATAGATG	ATCAACACTG	GCCGCGGATC	ACAACTGTTA
4921	TAATCATCGG	CATAGTATAT	CGGCATAGTA	TAATACGACT	CACTATAGGA	GGGCCACCAT
	ATTAGTAGCC	GTATCATATA	GCCGTATCAT	ATTATGCTGA	GTGATATCCT	CCCGGTGGTA
4981	GTCGACTACT	AACCTTCTTC	TCTTTCCTAC	AGCTGAGATC	ACCGGTAGGA	GGGCCATCAT
	CAGCTGATGA	TTGGAAGAAG	AGAAAGGATG	TCGACTCTAG	TGGCCATCCT	CCCGGTAGTA
5041	GAAAAAGCCT	GAACTCACCG	CGACGTCTGT	CGCGAAGTTT	CTGATCGAAA	AGTTCGACAG
	CTTTTTCGGA	CTTGAGTGGC	GCTGCAGACA	GCGCTTCAAA	GACTAGCTTT	TCAAGCTGTC
5101	CGTCTCCGAC	CTGATGCAGC	TCTCGGAGGG	CGAAGAATCT	CGTGCTTTCA	GCTTCGATGT
	CCACACCCTC.	CACTACCTCC	-NONGRETTECC	COTTOTTO	GCACGAAAGI	CGAAGCTACA
5161	AGGAGGGCGT	GGATATGTCC	TGCGGGTAAA	TAGCTGCGCC	GATGGTTTCT	ACAAAGATCG
	TCCTCCCGCA	CCTATACAGG	ACGCCCATTT	ATCGACGCGG	CTACCAAAGA	TGTTTCTAGC
5221	TTATGTTTAT	CGGCACTTTG	CATCGGCCGC	GCTCCCGATT	CCGGAAGTGC	TTGACATTGG
	AATACAAATA	GCCGTGAAAC	GTAGCCGGCG	CGAGGGCTAA	GGCCTTCACG	AACTGTAACC
5281	GGAATTCAGC	GAGAGCCTGA	CCTATTGCAT	CTCCCGCCGT	GCACAGGGTG	TCACGTTGCA
	CCTTAAGTCG	CTCTCGGACT	GGATAACGTA	GAGGGCGGCA	CGTGTCCCAC	AGTGCAACGT
5341	AGACCTGCCT	GAAACCGAAC	TGCCCGCTGT	TCTGCAACCC	GTCGCGGAGC	TCATGGATGC
	TCTGGACGGA	CTTTGGCTTG	ACGGGCGACA	AGACGTTGGG	CAGCGCCTCG	AGTACCTACG
5401	GATCGCTGCG	GCCGATCTTA	GCCAGACGAG	CGGGTTCGGC	CCATTCGGAC	CGCAAGGAAT
	CTAGCGACGC	CGGCTAGAAT	CGGTCTGCTC	GCCCAAGCCG	GGTAAGCCTG	GCGTTCCTTA
5461	CGGTCAATAC	ACTACATGGC	GTGATTTCAT	ATGCGCGATT	GCTGATCCCC	ATGTGTATCA
	GCCAGTTATG	TGATGTACCG	CACTAAAGTA	TACGCGCTAA	CGACTAGGGG	TACACATAGT
5521	CTGGCAAACT	GTGATGGACG	ACACCGTCAG	TGCGTCCGTC	GCGCAGGCTC	TCGATGAGCT
	GACCGTTTGA	CACTACCTGC	TGTGGCAGTC	ACGCAGGCAG	CGCGTCCGAG	AGCTACTCGA
5581	GATGCTTTGG	GCCGAGGACT	GCCCCGAAGT	CCGGCACCTC	GTGCACGCGG	ATTTCGGCTC
	CTACGAAACC	CGGCTCCTGA	CGGGGCTTCA	GGCCGTGGAG	CACGTGCGCC	TAAAGCCGAG
5641	CAACAATGTC	CTGACGGACA	ATGGCCGCAT	AACAGCGGTC	ATTGACTGGA	GCGAGGCGAT
	GTTGTTACAG	GACTGCCTGT	TACCGGCGTA	TTGTCGCCAG	TAACTGACCT	CGCTCCGCTA
5701	GTTCGGGGAT	TCCCAATACG	AGGTCGCCAA	CATCTTCTTC	TGGAGGCCGT	GGTTGGCTTG
	CAAGCCCCTA	AGGGTTATGC	TCCAGCGGTT	GTAGAAGAAC	ACCTCCGGCA	CCAACCGAAC
5761	TATGGAGCAG	CAGACGCGCT	ACTTCGAGCG	GAGGCATCCG	GAGCTTGCAG	GATCGCCGCG
E001	ATACCTCGTC	GTCTGCGCGA	TGAAGCTCGC	CTCCGTAGGC	CTCGAACGTC	CTAGCGGCGC
5821	GCTCCGGGCG	TATATGCTCC	GCATTGGTCT	TGACCAACTC	TATCAGAGCT	TGGTTGACGG
5881	CGAGGCCCGC	ATATACGAGG	CGTAACCAGA	ACTGGTTGAG	ATAGTCTCGA	ACCAACTGCC
2881	CAATTTCGAT	GATGCAGCTT	GGGCGCAGGG	TCGATGCGAC	GCAATCGTCC	GATCCGGAGC
5941	GTTAAAGCTA	CTACGTCGAA	CCCGCGTCCC	AGCTACGCTG	CGTTAGCAGG	CTAGGCCTCG
3941	CCCCTCACAC	GGGCGTACAC	AAATCGCCCG	CAGAAGCGCG	GCCGTCTGGA	CCGATGGCTG
6001	TCTACAACTC	CCCGCATGTG	TTTAGCGGGC	GTCTTCGCGC	CGGCAGACCT	GGCTACCGAC
0001	A CA TCTTCA C	GCGTCTGCGT	TCGACCAGGC	TGCGCGTTCT	CGCGGCCATA	GCAACCGACG
6061	TACCCCCCTTC	CGCAGACGCA	AGCTGGTCCG	ACGCGCAAGA	GCGCCGGTAT	CGTTGGCTGC
0001	ATCCCCCAAC	CGCCCTCGCC	GGCAGCAAGA	AGCCACGGAA	GTCCGCCCGG	AGCAGAAAAT
6121	CCCCACCCTA	GCGGGAGCGG	CCGTCGTTCT	TCGGTGCCTT	CAGGCGGGCC	TCGTCTTTTA
0121	CCCCACGCIA	CTGCGGGTTT	ATATAGACGG	TCCCCACGGG	ATGGGGAAAA	CCACCACCAC
	CGGGIGCGAT	GACGCCCAAA	TATATCTGCC	AGGGGTGCCC	TACCCCTTTT	GGTGGTGGTG

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FIG. 1E





6181		CTCCCCCCTCC	CERTOCOCOCO A	CC3 M3 MCCMC	<b>T1</b> COT1 COCC	, aaaa, ma, a
0101		GTGGCCCTGG				
C243		CACCGGGACC				
6241		GTGCTGGGGG				
		CACGACCCCC				
6301		GGTGAGATAT				
		CCACTCTATA				
6361		ATGCCTTATG			-	
		TACGGAATAC			<del>-</del>	
6421		TCACATGCCC				
		AGTGTACGGG				
6481		CTGTGCTACC				
		<del>CACACCATES</del>				
6541		TTCGTGGCCC				
		AAGCACCGGG				
6601		GAGGACAGAC				
	CCGGGAAGGC	CTCCTGTCTG	TGTAGCTGGC	GGACCGGTTT	GCGGTCGCGG	GGCCGCTCGC
6661	GCTGGACCTG	GCTATGCTGG	CTGCGATTCG	CCGCGTTTAC	GGGCTACTTG	CCAATACGGT
		CGATACGACC				
6721		CAGTGCGGCG				
	CGCCATAGAC	GTCACGCCGC	CCAGCACCGC	CCTCCTGACC	CCTGTCGAAA	GCCCCTGCCG
6781	CGTGCCGCCC	CAGGGTGCCG	AGCCCCAGAG	CAACGCGGGC	CCACGACCCC	ATACGGGGGA
		GTCCCACGGC				
6841	CACGTTATTT	ACCCTGTTTC	GGGCCCCCGA	GTTGCTGGCC	CCCAACGGCG	ACCTGTATAA
	GTGCAATAAA	TGGGACAAAG	CCCGGGGGCT	CAACGACCGG	GGGTTGCCGC	TGGACATATT
6901	CGTGTTTGCC	TGGGCCTTGG	ACGTCTTGGC	CAAACGCCTC	CGTTCCATGC	ACGTCTTTAT
	GCACAAACGG	ACCCGGAACC	TGCAGAACCG	GTTTGCGGAG	GCAAGGTACG	TGCAGAAATA
6961	CCTGGATTAC	GACCAATCGC	CCGCCGGCTG	CCGGGACGCC	CTGCTGCAAC	TTACCTCCGG
	GGACCTAATG	CTGGTTAGCG	GGCGGCCGAC	GGCCCTGCGG	GACGACGTTG	AATGGAGGCC
7021	GATGGTCCAG	ACCCACGTCA	CCACCCCGG	CTCCATACCG	ACGATATGCG	ACCTGGCGCG
	CTACCAGGTC	TGGGTGCAGT	GGTGGGGGCC	GAGGTATGGC	TGCTATACGC	TGGACCGCGC
7081	CACGTTTGCC	CGGGAGATGG	GGGAGGCTAA	CTGAGTCGAG	AATTCGCTAG	AGGGCCCTAT
	GTGCAAACGG	GCCCTCTACC	CCCTCCGATT	GACTCAGCTC	TTAAGCGATC	TCCCGGGATA
7141	TCTATAGTGT	CACCTAAATG	CTAGAGCTCG	CTGATCAGCC	TCGACTGTGC	CTTCTAGTTG
	AGATATCACA	GTGGATTTAC	GATCTCGAGC	GACTAGTCGG	AGCTGACACG	GAAGATCAAC
7201	CCAGCCATCT	GTTGTTTGCC	CCTCCCCGT	GCCTTCCTTG	ACCCTGGAAG	GTGCCACTCC
	GGTCGGTAGA	CAACAAACGG	GGAGGGGGCA	CGGAAGGAAC	TGGGACCTTC	CACGGTGAGG
7261	CACTGTCCTT	TCCTAATAAA	ATGAGGAAAT	TGCATCGCAT	TGTCTGAGTA	GGTGTCATTC
	GTGACAGGAA	AGGATTATTT	TACTCCTTTA	ACGTAGCGTA	ACAGACTCAT	CCACAGTAAG
7321	TATTCTGGGG	GGTGGGGTGG	GGCAGGACAG	CAAGGGGGAG	GATTGGGAAG	ACAATAGCAG
	ATAAGACCCC	CCACCCCACC	CCGTCCTGTC	GTTCCCCCTC	CTAACCCTTC	TGTTATCGTC
7381	GCATGCGCAG	GGCCCAATTG	CTCGAGCGGC	CGCAATAAAA	TATCTTTATT	TTCATTACAT
	CGTACGCGTC	CCGGGTTAAC	GAGCTCGCCG	GCGTTATTTT	ATAGAAATAA	AAGTAATGTA
7441	CTGTGTGTTG	GTTTTTTGTG	TGAATCGTAA	CTAACATACG	CTCTCCATCA	AAACAAAACG
	GACACACAAC	CAAAAAACAC	ACTTAGCATT	GATTGTATGC	GAGAGGTAGT	TTTGTTTTGC
7501	AAACAAAACA	AACTAGCAAA	ATAGGCTGTC	CCCAGTGCAA	GTGCAGGTGC	CAGAACATTT
	TTTGTTTTGT	TTGATCGTTT	TATCCGACAG	GGGTCACGTT	CACGTCCACG	GTCTTGTAAA

FIG. 1F

Sheet 7 of 7
Title: SELECTION SYSTEMS FOR GENETICALLY MODIFIED

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CTCTATCGAA GGATCTGCGA TCGCTCCGGT GCCCGTCAGT GGGCAGAGCG CACATCGCCC 7561 GAGATAGCTT CCTAGACGCT AGCGAGGCCA CGGGCAGTCA CCCGTCTCGC GTGTAGCGGG ACAGTCCCCG AGAAGTTGGG GGGAGGGGTC GGCAATTGAA CCGGTGCCTA GAGAAGGTGG 7621 TGTCAGGGGC TCTTCAACCC CCCTCCCCAG CCGTTAACTT GGCCACGGAT CTCTTCCACC CGCGGGGTAA ACTGGGAAAG TGATGTCGTG TACTGGCTCC GCCTTTTTCC CGAGGGTGGG 7681 GCGCCCCATT TGACCCTTTC ACTACAGCAC ATGACCGAGG CGGAAAAAGG GCTCCCACCC GGAGAACCGT ATATAAGTGC AGTAGTCGCC GTGAACGTTC TTTTTCGCAA CGGGTTTGCC 7741 CCTCTTGGCA TATATTCACG TCATCAGCGG CACTTGCAAG AAAAAGCGTT GCCCAAACGG Ø1 7801 GCCAGAACAC AGCTGAAGCT TCGAGGGGCT CGCATCTCTC CTTCACGCGC CCGCCGCCCT CGGTCTTGTG TCGACTTCGA AGCTCCCCGA GCGTAGAGAG GAAGTGCGCG GGCGGCGGGA ACCTGAGGCC GCCATCCACG CCGGTTGAGT CGCGTTCTGC CGCCTCCCGC CTGTGGTGCC 7861 æ TGGACTCCGG CGGTAGGTGC GGCCAACTCA GCGCAAGACG GCGGAGGGCG GACACCACGG TCCTGAACTG CGTCCGCCGT CTAGGTAAGT TTAAAGCTCA GGTCGAGACC GGGCCTTTGT 7921 AGGACTTGAC GCAGGCGGCA GATCCATTCA AATTTCGAGT CCAGCTCTGG CCCGGAAACA Ŋ CCGGCGCTCC CTTGGAGCCT ACCTAGACTC AGCCGGCTCT CCACGCTTTG CCTGACCCTG 7981 GGCCGCGAGG GAACCTCGGA TGGATCTGAG TCGGCCGAGA GGTGCGAAAC GGACTGGGAC CTTGCTCAAC TCTACGTCTT TGTTTCGTTT TCTGTTCTGC GCCGTTACAG ATCCAAGCTG GAACGAGTTG AGATGCAGAA ACAAAGCAAA AGACAAGACG CGGCAATGTC TAGGTTCGAC TGACCGGCGC CTACGTAAGT GATATCTACT AGATTTATCA AAAAGAGTGT TGACTTCTGA 8101 ACTGGCCGCG GATGCATTCA CTATAGATGA TCTAAATAGT TTTTCTCACA ACTGAACACT GCGCTCACAA TTGATACTTA GATTCATCGA GAGGGACACG TCGACTACTA ACCTTCTTCT 8161 CGCGAGTGTT AACTATGAAT CTAAGTAGCT CTCCCTGTGC AGCTGATGAT TGGAAGAAGA CTTTCCTACA GCTGAGAT 8221 GAAAGGATGT CGACTCTA

FIG. 1G